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GCGCAAGCCCTGGGCCCCGGAGGTGCAGGAGGAGGAGGAGCCCCCAGGGGGGACGAGGACGTGCTGTTCAGGGTGACAAA

FIG. 1A

0 8	160	240	320	400	480
ATGGATTTCGGACTGGCCTCCTGCGGGGCTTCTGGGGCTCCTCGGCCAGTCCCTCCAGGTGAAGCCCCTGCA	GGTGGAGCCCCCGGAGCCGTGGCCGTGGCCTTGGGCGCCTCGCCCAGCTCACCTGCCGCCTGCGCGCGC	GCGGGGCCTCGGTGCAGTGGCGGGGCCTGGGCGCGCGGGCGAGTCGGACACGGGCCGCAGCGTCCTCACC R G A S V Q W R G L D T S L G A V Q S D T G R S V L T	GTGCGCAACGCCTCGCTGTCGGCGGCCCGGGACCCGCGTGGGGCTCCTGCGGGGGCCGCACCTTCCAGCACACCGT V R N A S L S A A G T R V C V G S C G G R T F Q H T V	GCAGCTCCTTGTGTACGCCTTCCCGGACCAGCTGACCGTCTCCCCAGCAGCCCTGGTGCCTGGTGACCCGGAGGTGGCCT Q L L V Y A F P D Q L T V S P A A L V P G D P E V A	GTACGGCCCACAAAGTCACGCCCCGTGGACCCCAGGCGCCTCCTTCTCCCTGCTCGGGGGCCAGGAACTGGAGGGG C ta hkvtpvdprocessessessessessessessessessessessessess

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	CAC 800	3 S	S S	GTG 1040	scrg 1120 A	CAG 1200
SECOCOCOCOCOCOCOCOCOCOCOCOCOCOCOCOCOCOCO	CCCAACACCACCTCCCGGAGTCTCCCGACACCTCCCGGAGTCTCCCGACACCACCTCCCAGGAGCCTCCCGACACACAC	CACCTCCCGGAGCCTCCCGACACCTCCCAGGAGCCTCCCGACACCTCCCGGAGCCTCCCGACAAGACCTCCC T S Q E P D T T S Q E P D T T S P E P D K T S	CGGAGCCCGCCCCCAGGGCTCCACACACCCCCAGGAGCCCAGGCTCCACCAGGACTCGCCGCCCTGAGATCTCC P E P A P Q Q G S T H T P R S P G S T R T R P E I S	CAGGCTGGGCCCACGCAGGAAGTGATCCCAACAGGCTCGTCCAAACCTGCGGGTGACCAGCTGCCGGGGTCTGTG Q A G P T Q G E V I P T G S S K P A G D Q L P A L W	GACCAGCAGTGCGGTGCTGCTGCTCCTGGCCTTGCCCACGTATCACCTCTGGAAACGCTGCCGGCACCTGGCTG TSSAVLGLLLALPTYHLWKRCRHLAALA	AGGACGACCCACCCACCAGCTTCTCTGAGGCTTCTGCCCCAGGTGTCGGCCTGGGCTTAAGGGGGACCGGCCAG

0.0	40	0	3/27
1360	1440	1520	1600
GGTCAGGGCAAACCTGCCTCCCATTCTACTCAAAGTCATCCCTCTGCTCACAGAGATGGATG	TTGGAGAAGCTCATCAGAAACTCAAAAGAAGGCCACTGTTTGTCTCACCTACCCATGACCTGAAGCCCCTTCCTT	TCCCCACCTTTCTGGACGGAACCACGTACTTTTACATACA	CAAGCTGTGCCCTGACCCTGGGCCCCTGTCGTCAGGACCTCCTGAGGCTTTGGCCAAATAAACCTCCTAAAATGATAA

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GCGCAAGCCCTGGGCCCGGAGGAGGAGGAGCCCCCAGGGGGGACGACGTGCTGTTCAGGGTGACAGA

FIG. 2A

2A 80	3C 160	3C 240	320 v	T 400	iG 480
ATGGATTTCGGACTGGCCTCCTGCGGGCTTCTGGGGCTCCTCCTCGGCCAGTCCCTCCAGGTGAAGCCCCTGCA	GGTGGAGCCCCCGGAGCCGTGGCCGTGGCCTTGGGCGCCTCGCCCAGCTCACCTGCCGCCTGCCGCGGGCCCCCCCC	GCGGGGCCTCGGTGCCGGGGCCTGGACACCAGCCTGGGCGGCGGGCG	GTGCGCAACGCCTGTCGGCGGCCGGGACCCGCGTGTGCGTGGGCTCCTGCGGGGGCCGCACCTTCCAGCACACGT ${ m V}$ R ${ m N}$ A ${ m S}$ L ${ m S}$ A ${ m A}$ G ${ m IC}$ V ${ m G}$ V ${ m G}$ O ${ m G}$ C ${ m G}$ G ${ m G}$ G ${ m G}$ C ${ m G}$ G ${ m G}$ R T F ${ m Q}$ H T V	GCAGCTCCTTGTGTACGCCTTCCCGGACCAGCTGACCGTGCTGCTGGTGACCCGGAGGTGGCCT Q L L V T A F P D Q L T V S P A A L V P G D P E V A	GTACGGCCCACAAAGTCACGCCCGTGGACCCCCAACGCGCTCTCCTTCTCCCTGCTCGGGGGCCCAGGAACTGGAGGGG $\overline{\mathbb{C}}$ T P V D P N A L S F S L L V G G \mathbb{G} \mathbb{C} E L \mathbb{E} G
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FIG. 2C

CCGCCCAGGTCGGGATCAGCCCTCCTGAGTGGCCAGCCTTTCCCCCTGTGAAAGCAAATAGCTTGGACCCCTTCAAGT F G Q V G I S P S FGAGAACTGGTCAGGCCAAACTCCATTCTACTCAAAGTCATCCTTGTTCACAGAGATGGATG	1200
ITGCCTCTTTGGAGAAGCTCAGAAACTCAAAAGAAGGCCACTGTTTGTCTCACCTACCCATGACCTGAAGCCCCTCC	1360
CTGAGTGGTCCCCACCTTTCTGGACGGAACCACGTACTTTTTACATACA	1440
CGTAAGACCAAGCTGTGCCCTGACCACCCTGGGCCCCTGTCGTCAGGACCTCCTGAGGCTTTGGCAAATAAACCTCCTAA	27 1250

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80	160	240	320	400	480	560
AGCATGGATCGGGGCCTCCTCCTGCTGGCGGGCTTCTGGGGCTCCTCC	GAAGCCCCTGCAGGTGGAGCCGGTGGTGGCCGTGGCCCTGGGCCCTTTCGCCAGCTCACCTGCCGCCTGG	ACTGCGCGGACCGGGGCCACGGTGCAGTGGGGGCGTGGGGCGCGGGGGGGG	AGCGTCCTCACCGTGCGCCTCGCTGTCGGCGGCCGGGACCCGTGTGTGT	CCAGCACCACCGTGCGGTCCTTGCCGGACCAGCTGACCATCTCCCCGGCAGCCCTGGTGCCTGGTGACC Q H T V R L L V Y A F P D Q L T I S P A A L V P G D	CGGAGGTGGCCTGTACGGCCCACAAGTCACGCCTGTGGACCCCCAATGCGCTCTTCTCCCTGCTCCTGGGGGGACCAG	GAACTGGAGGGGCCCCAGGCCCCGGAGGTGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG

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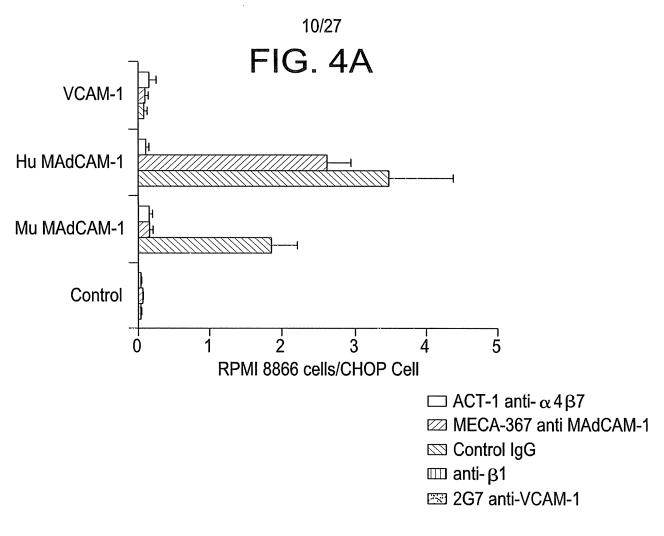
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640	720	800	880	960
CAGGGTGACAGACGCTGCCGACCCTGGCAACCCCTGTCCTGCCGCGCTCTACTGCCAGGCCACGATGAGGC R $ m V$ I $ m E$ R $ m W$ R $ m I$ P $ m I$ L A T P $ m V$ I P A L Y $ m C$ Q A T M R	TGCCTGGCTTGGAGCTCAGCCCACCGCCATCCCGGTCCTGCACGGCCCGGGAGCCCCCCGACCCCCGACACGACCTCCCGGGAGCCCCCCCGACACGACC	TCCCCGGAACCCCGGGCCGCGAGCCCCCCCCCCCGGGGCTCCACACACCCCCAGGGGCTCCACACACCACCAGGGCTCCACACACCACCAGGGCTCCACACACCACCAGGGCTCCACACCACCAGGGCTCCACACCACCAGGGCTCCACACCACCAGGGCTCCACACCACCAGGGCTCCACACCACCAGGGCTCCACACCACCAGGGCTCCACAGGGCTCCACAGGGCTCCACAGGGCTCCACAGGGCTCCACAGGGCTCCACAGGGCTCCACAGGGCTCCACGGGGCTCCACGGGGCTCCACGGGGCTCCACGGGGCTCCAGGGGCTCCACGGGGCTCCAGGGGCTCCACGGGGCTCCACGGGGCTCCACGGGGCTCCAGGGCTCCACGGGGCTCCAGGGGCCTCAGGGGCCTCAGGGGCCTCAGGGGCCTCAGGGGCCTCAGGGGCCTCAGGGGCCTCAGGGGCCTCAGGGGCCTCAGGGGCCTCAGGGGCCTCAGGGGCCTCAGGGGCCTCAGGGGCCTCCAGGGGCCTCAGGGGCCTCCAGGGGCCTCAGGGCCTCAGGGGCCTCCAGGGGCCTCCAGGGGCCTCCAGGGGCCTCCAGGGGCCTCCAGGGGCCTCCAGGGGCCCCAGGGGCCTCCAGGGGCCCCCAGGGGCCTCCAGGGGCCTCCAGGGGCCTCCAGGGGCCTCCAGGGGCCTCCAGGGGCCTCCAGGGGCCTCCAGGGGCCCCCCAGGGGCCTCCAGGGGCCCCCCAGGGGCCCCCCAGGGGCCTCCAGGGGCCCCCCAGGGGCCCCCCCAGGGGCCCCCCCAGGGGCCCCCC	TACCAGGACTTGCCGCCCTGAGATCTCCCAGGCTGGGCCCCACGGGAGAGTGATCCCAACAGGCTCGTCCAAACCTA	CGGGTGACCAGCTGCCCGCGGCTCTGTGGACCAGCAGTGCGGTGCTGGGACTGCTGCTCCTGGCTTTGCCCCACCTACCAC

AAGGGAAAATAGGTTGGACCCCTTCAAGCTGAGAACTGGTCGGGGCAAACCTGCCTCCCATTCTATTCAAAGTCATCGCT

FIG. 3C

1200	1280	1360	9/27 9/27	1520	1600	1680
CTGGTCACAGAGGGACGCACATTCTGATTGCCTCCTTTGGAAAGGCTCATCAGAAACTCAAAAGGAGGTGATCGTTTG	TCCCGCCTACCCGTGACCTGGAAGCCCCCCCCCCCCGCTCGAGTGACCCTTGACTTTCTGGACGGAACCAACGTACTTCTTA	CATATATTGATTGATGTGTCATATCTCCCTAAAATGCGTAAAACCAGCTGTGCCCCGACCACCTTGGGCCCCTGCCATCA	GGACCTCCTGAGGCTTTGGCAAATAAACCTCCTAAAAGGATAGAAACTGAAACTTGTGGCCGGGCGCGGGTGGCTCAAGCC	TGTAATCCCAGCACTTTGGGAGGCCGAGGTGGGTCACGAGGTCAGGAGATCGAGACCATCCTGGCTAACCCGTGAA	ACCCCGTCTCTACTAAAAATACAAAATTAGCCGGGAGCGGTGGCGGGGGCGCCTGTAGTCCCCAGCTACTCGGGAGGCTG	AGGCAGGAGAATGGCGTGAACCCGGGAGGCGGGGGCTTGCAGTGAGCTGAGATCCGGCCCACTGCACTCCAGCCTGGGGGAC



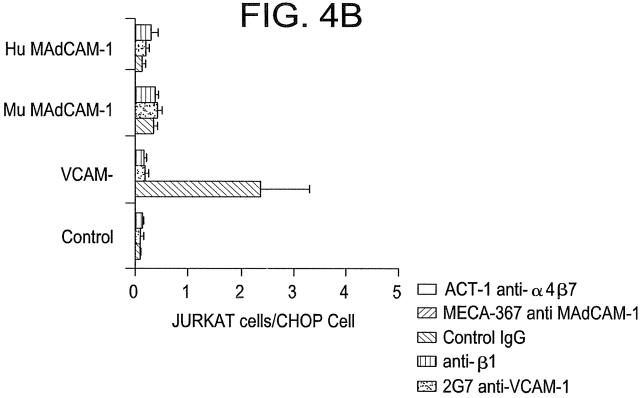
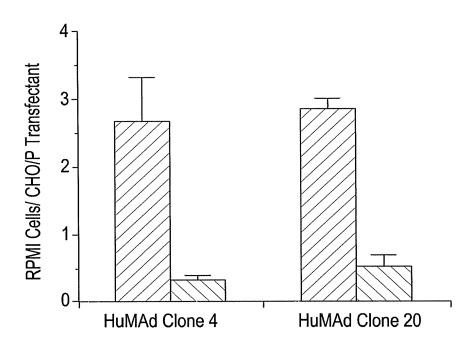


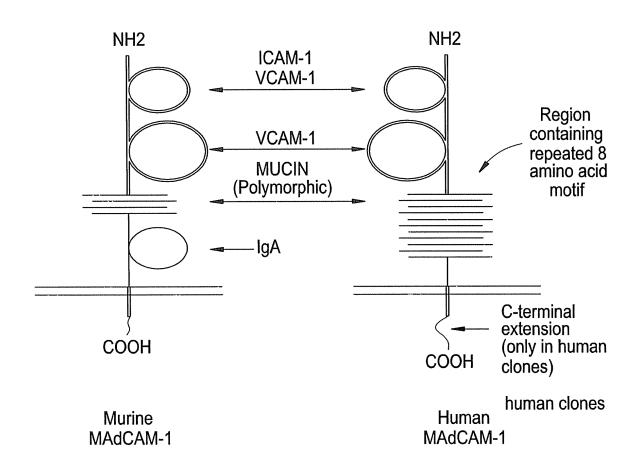
FIG. 5



∠ Control IgG

 \triangle Act-1 anti- α 4 β 7

FIG. 6



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FIG. 7B

Structural Damage

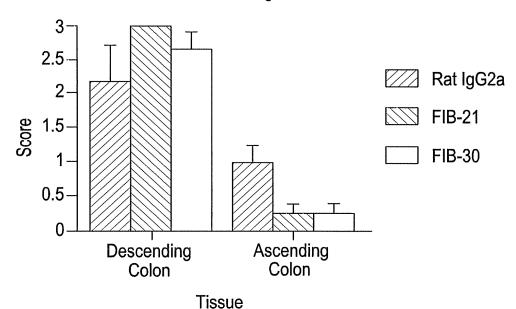
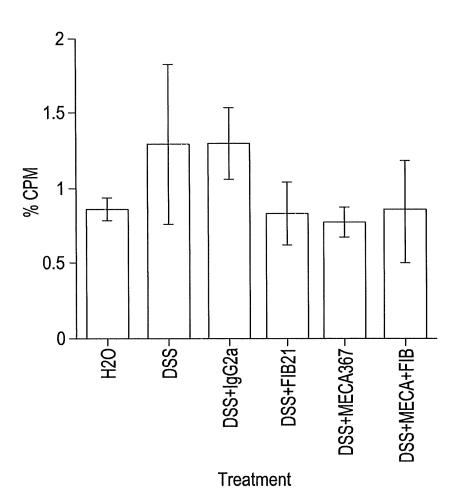


FIG. 8



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FIG. 9

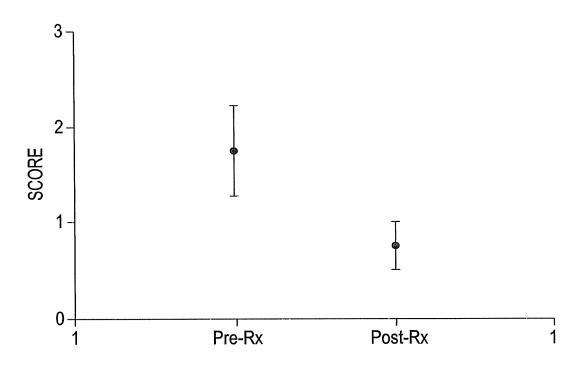


FIG. 10

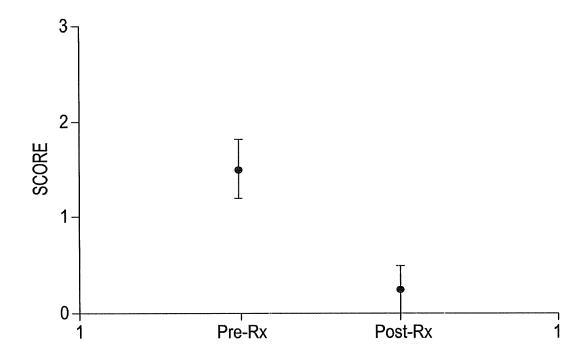


FIG. 11

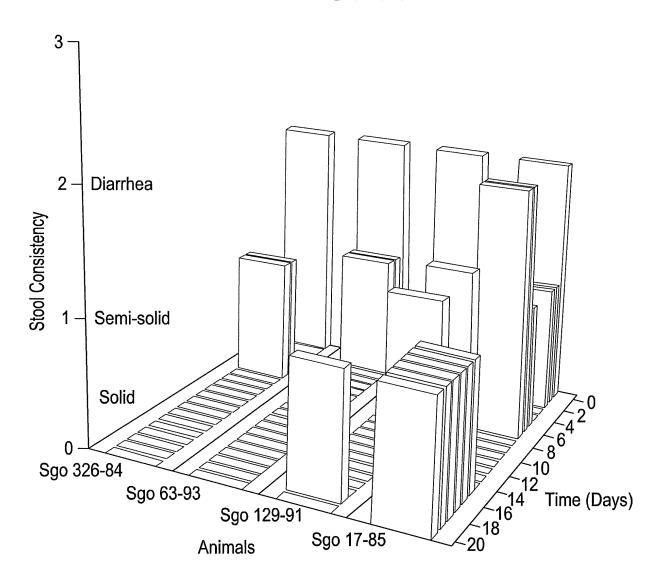


FIG. 12

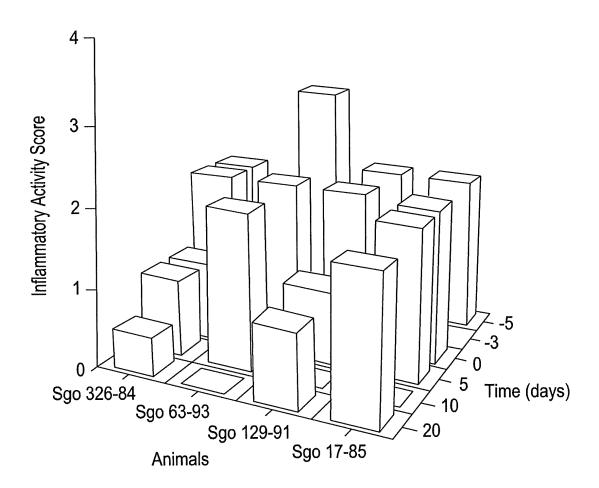


FIG. 13

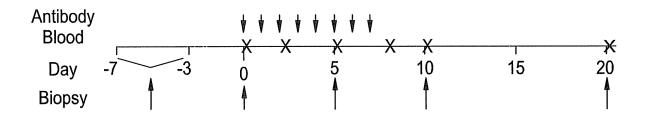
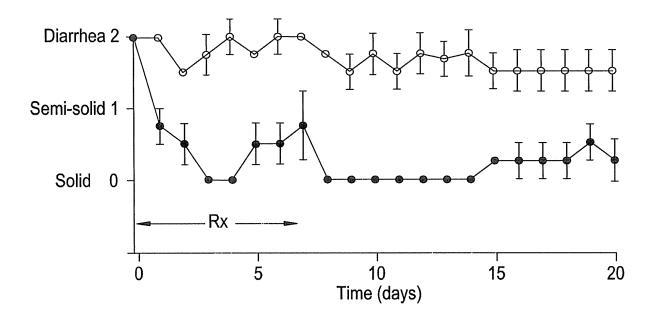


FIG. 14

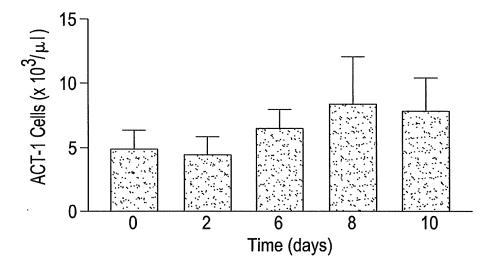


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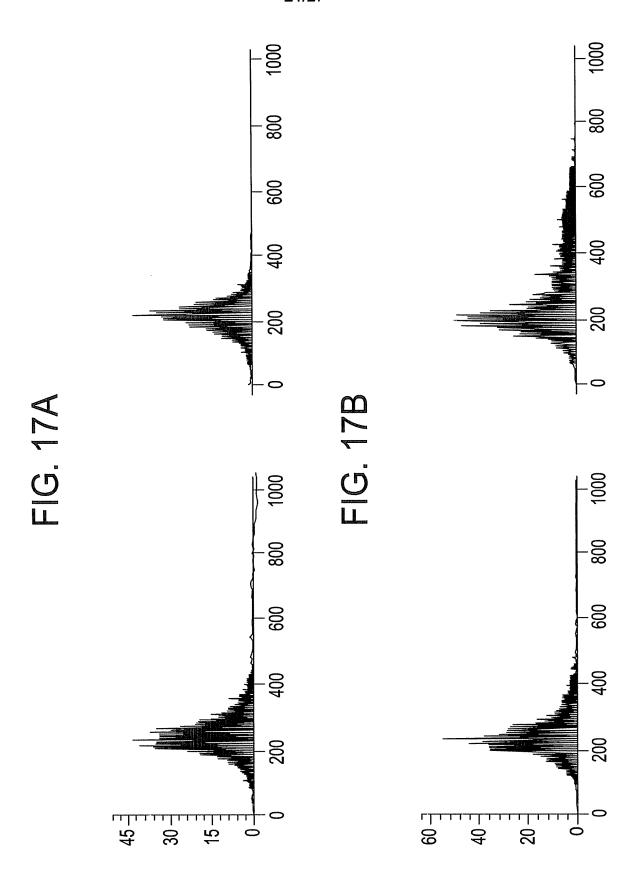
FIG. 15

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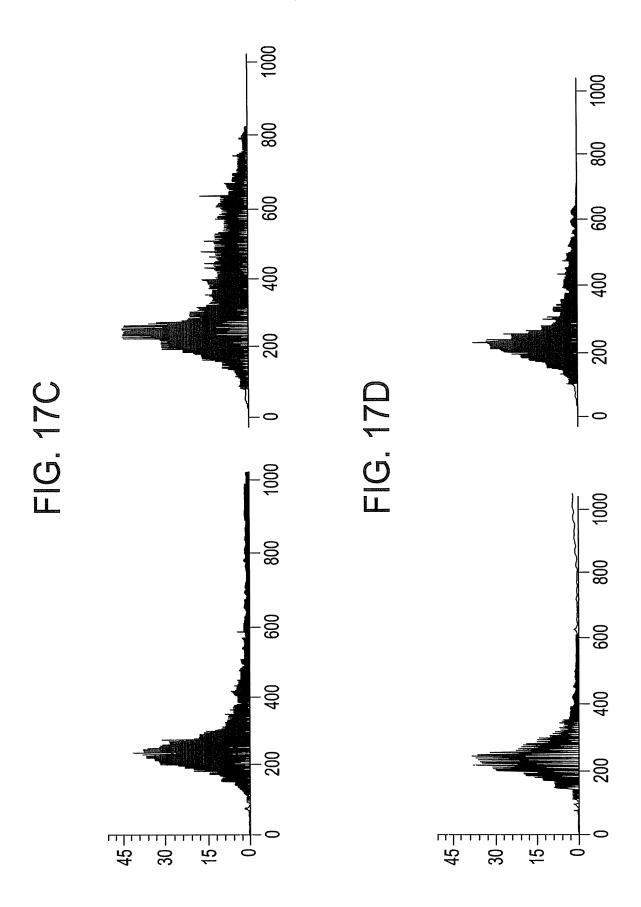
FIG. 16



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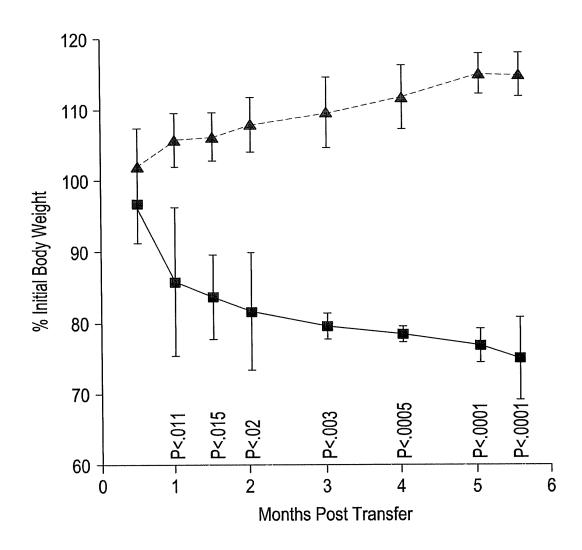
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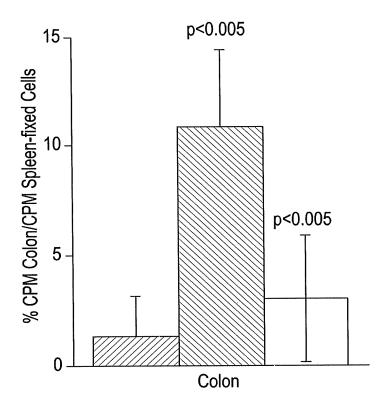


FIG. 18



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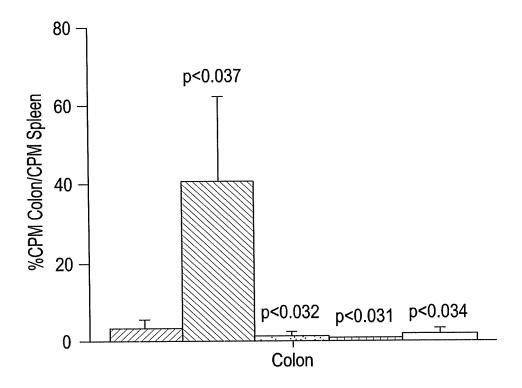
FIG. 19



- CD45RB-lo/Rat lgG2a
- CD45RB-hi/Rat IgG2a
- CD45RB-hi/FIB-504+MECA-367

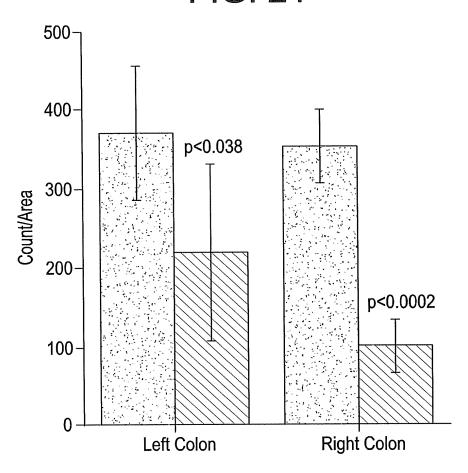
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FIG. 20



- CD45RBhi/FIB-504
- CD45RBhi/MECA-367
- CD45RBhi/FIB504+MECA367

FIG. 21



Rat IgG2a

FIB504+MECA367